

## SEQLIST.TXT

## SEQUENCE LISTING

&lt;110&gt; Albert, Lai

&lt;120&gt; NOVEL SPLICING VARIANTS OF HUMAN DKK11

&lt;130&gt; PP023359.0003

&lt;140&gt; 10/574182

&lt;141&gt; 2007-05-31

&lt;150&gt; PCT/US04/34256

&lt;151&gt; 2004-09-30

&lt;150&gt; 60/507682

&lt;151&gt; 2003-09-30

&lt;160&gt; 26

&lt;170&gt; FastSEQ for Windows Version 4.0

&lt;210&gt; 1

&lt;211&gt; 819

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 1

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agtccttgg gtctcacagg cctccagagc ctactccaag gcttcagccg acttttcctg 180  
aaaggtaacc tgcttcgggg catagacagc ttattctctg ccccatgga cttccggggc 240  
ctccctggga actaccacaa agaggagaac caggagcacc agctggggaa caacaccctc 300  
tccaggacc tccagatcga caagaggacc gacaacaaga caggagaggt gctgatctcc 360  
gagaatgtgg tggcatccat tcaaccagcg gaggggagct tcgagggtga tttgaaggt 420  
cccagatgg aggagaaga ggccttggta cccatccaga aggccacgga cagcttccac 480  
acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggcg gaggtccac 540  
caggatgccc tggagggcgg ccactggctc agcgagaagc gacaccgcct gcaggccatc 600  
cgggatggac tccgcaaggg gaccacaag gacgtcctag aagagggac cgagactcc 660  
tcccactcca ggctgtcccc ccgaaagacc cacttactgt acatcctcag gcccctctcg 720  
cagctgttagg ggtggggacc ggggagcacc tgcctgtacccc cccatcaga ccctgcccc 780  
agcaccatat gaaaataaaag ttctttctta catctaaca 819

&lt;210&gt; 2

&lt;211&gt; 242

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 2

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
1 5 10 15  
Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
20 25 30  
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
35 40 45  
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
50 55 60  
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
65 70 75 80  
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
85 90 95  
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Arg Thr Asp Asn

## SEQLIST.TXT

100	105	110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln		
115	120	125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu		
130	135	140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His		
145	150	155
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg		
165	170	175
Arg Arg Ser His Gln Asp Ala Leu Glu Gly His Trp Leu Ser Glu		
180	185	190
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr		
195	200	205
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser His Ser Arg		
210	215	220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg		
225	230	235
Gln Leu		240

&lt;210&gt; 3

&lt;211&gt; 733

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 3

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gctccctct	accctggta	tcccctccgc	tgtagctct	atccatgtat	ctgacgcccc	120
agagagctcc	ttgggtctca	caggcctcca	gaggctactc	caaggcttca	gccgactttt	180
cctgaaagggt	aacctgcttc	ggggcataga	cagcttattc	tctgccccca	tggacttccg	240
gggcctccct	gggaactacc	acaagagga	gaaccaggag	caccagctgg	ggaacaacac	300
cctctccagc	cacccctcaga	tcgacaagat	gaccgacaaac	aagacaggag	aggtgctgat	360
ctccgagaat	gtgggtggcat	ccattcaacc	agcggaggggg	agtttcgagg	tgatttgaa	420
gttaccagg	atggaggaga	aggaggccat	gttaccatcc	cagaaggcca	cggacagctt	480
ccacacagaa	ctccatcccc	gggtggccctt	ctggatcatc	aagctgcccac	ggcggagggtc	540
ccaccaggat	gccctggagg	gcccactg	gctcagcgag	aagcgacacc	gcctgcaggc	600
catccggat	gactccgca	aggggaccca	caaggacgtc	ctagaagagg	ggaccgagag	660
ctcccccac	tccaggctgt	ccccccgaaa	gaccactta	ctgtacatcc	tcaggccctc	720
tcggcagctg	tag					733

&lt;210&gt; 4

&lt;211&gt; 242

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 4

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val			
1	5	10	15
Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro			
20	25	30	
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu			
35	40	45	
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu			
50	55	60	
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly			
65	70	75	80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly			
85	90	95	
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn			
100	105	110	
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln			
115	120	125	

## SEQLIST.TXT

Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu  
 130 135 140  
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His  
 145 150 155 160  
 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg  
 165 170 175  
 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu  
 180 185 190  
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr  
 195 200 205  
 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser His Ser Arg  
 210 215 220  
 Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg  
 225 230 235 240  
 Gln Leu

&lt;210&gt; 5

&lt;211&gt; 733

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 5

caccatggga gaagcctccc cacctgcccc cgcaaggcgg catctgctgg tcctgctgct 60  
 gtcctctctt accctggta tcccctccac tgcaagcttccatccatgtatgt ctgacgccc 120  
 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcctca gcccactttt 180  
 cctgaaaggt aacctgcttc gggccataga cagcttattc tctgccccca tggacttccg 240  
 gggcctccctt gggactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
 cctctccagc cacctccaga tcgacaagat gaccgacaac aagacaggag aggtgctgat 360  
 ctccgagaat gtgggtggcat ccattcaacc agcggagggg agcttcgagg gtgattt 420  
 ggtacccagg atggaggaga aggaggccct ggtacccatc cagaaggcca cggacagctt 480  
 ccacacagaa ctccatcccc gggggccctt ctggatcatt aagctgcccac ggcggagg 540  
 ccaccaggat gcccctggagg gcccactg gctcagcgg aagcgacacc gcctgcaggc 600  
 catccggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660  
 ctccctccac tccaggctgt ccccccggaaa gaccctta ctgtacatcc tcaggccctc 720  
 tcggcagctg tag 733

&lt;210&gt; 6

&lt;211&gt; 242

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 6

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
 50 55 60  
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
 65 70 75 80  
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
 85 90 95  
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn  
 100 105 110  
 Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln  
 115 120 125  
 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu  
 130 135 140  
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His

## SEQLIST.TXT

145	150	155	160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg			
165	170	175	
Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu			
180	185	190	
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr			
195	200	205	
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser His Ser Arg			
210	215	220	
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg			
225	230	235	240
Gln Leu			

<210> 7  
<211> 733  
<212> DNA  
<213> homo sapien

<400> 7  
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gctccctctt accctggta tccccctccac tgcatctt atccatgtat ctgacgcccc 120  
agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180  
cctgaaaggt aacctgcttc gggcataga cagcttattc tctgccccca tggacttccg 240  
gggcctccct gggactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
cctctccagc cacctccaga tcgacaagat gaccgacaa aagacaggag aggtgctgat 360  
ctccgagaat gtgggtggcat ccattcaacc agcggaggg agttcgagg gtgatttcaa 420  
ggtaccagg atggaggaga aggaggccct ggtaccatc cagaaggcca cggacagctt 480  
ccacacagaa ctccatcccc gggggccctt ctggatcatt aagctgcccac ggcggaggtc 540  
ccaccaggat gcccggagg gggccactg gctcagcgg aagcgacacc gcctgcaggc 600  
catccggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660  
ctccctccac tccaggctgt ccccccggaaa gacccactta ctgtacatcc tcaggccctc 720  
tcggcagctg tag 733

<210> 8  
<211> 242  
<212> PRT  
<213> homo sapien

<400> 8  
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
1 5 10 15  
Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro  
20 25 30  
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
35 40 45  
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
50 55 60  
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
65 70 75 80  
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
85 90 95  
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn  
100 105 110  
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln  
115 120 125  
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu  
130 135 140  
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His  
145 150 155 160  
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg  
165 170 175

## SEQLIST.TXT

Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu  
 180 185 190  
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr  
 195 200 205  
 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser His Ser Arg  
 210 215 220  
 Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg  
 225 230 235 240  
 Gln Leu

<210> 9  
 <211> 733  
 <212> DNA  
 <213> homo sapien

<400> 9  
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 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gcccacttt 180  
 cctgaaaggta accctgcttc ggggcataga cagtttattc tctgccccca tggacttccg 240  
 gggcctccctt gggaaactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
 cctctccagc caccctccaga tcgacaagat gaccgacaaac aagacaggag aggtgctgat 360  
 ctccgagaat gtgggtggcat ccattcaacc agcggagggg agcttcgagg gtgattgaa 420  
 ggtacccagg atggaggaga aggaggccctt ggtacccatc cagaaggcca cggacagctt 480  
 ccacacagaa ctccatcccc gggtggcattt ctggatcatt aagctgcccac ggcggaggc 540  
 ccaccaggat gcccctggagg gcagccactg gtcagcgg aagcgacacc gcctgcaggc 600  
 catccggat ggactccgca aggggaccca caaggacgtc ctaaaagagg ggaccgagag 660  
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 tcggcagctg tag 733

<210> 10  
 <211> 242  
 <212> PRT  
 <213> homo sapien

<400> 10  
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
 50 55 60  
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
 65 70 75 80  
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
 85 90 95  
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn  
 100 105 110  
 Lys Thr Gly Glu Val Leu Ile Ser Gln Glu Asn Val Val Ala Ser Ile Gln  
 115 120 125  
 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu  
 130 135 140  
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His  
 145 150 155 160  
 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg  
 165 170 175  
 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu Ser Glu  
 180 185 190  
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr

## SEQLIST.TXT

195	200	205
His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser His Ser Arg		
210	215	220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg		
225	230	235
Gln Leu		240

<210> 11  
 <211> 733  
 <212> DNA  
 <213> homo sapien

<400> 11  
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 gtcctctct accctggtga tccccctccac tgcagctcct atccatgtat ctgacgccc 120  
 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180  
 cctgaaagggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240  
 gggcctccct gggaaactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
 cctctccagc cacctccaga tcgacaagat gaccgacaac aagacaggag aggtgctgat 360  
 ctccgagaat gtggtggcat ccatcaacc agcggagggg agcttcgagg gtgatttcaa 420  
 ggtacccagg atggaggaga aggaggccct ggtacccatc cagaaggcca cggacagctt 480  
 ccacacagaa ctccatcccc gggggccctt ctggatcatt aagctgcccac ggcggaggc 540  
 ccaccaggat gcccctggagg gcccccaactg gctcagcgg aagcgacacc gcctgcaggc 600  
 catccggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660  
 ctccctccac tccaggctgt ccccccggaaa gaccaccca ctgtacatcc tcaggccctc 720  
 tcggcagctg tag 733

<210> 12  
 <211> 242  
 <212> PRT  
 <213> homo sapien

<400> 12  
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
 50 55 60  
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
 65 70 75 80  
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
 85 90 95  
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn  
 100 105 110  
 Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln  
 115 120 125  
 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu  
 130 135 140  
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His  
 145 150 155 160  
 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg  
 165 170 175  
 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu  
 180 185 190  
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr  
 195 200 205  
 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg  
 210 215 220

## SEQLIST.TXT

Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg  
 225 230 235 240  
 Gln Leu

&lt;210&gt; 13

&lt;211&gt; 640

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 13

caccatggga gaaggcctccc cacctgcccc cgcaaggcgg catctgctgg tcctgctgct 60  
 gctccctctt accctggta tcccctccgc tgcaagcttctt atccatgtg ctgacgccc 120  
 agagagctcc ttgggtctca caggcctcca gaggctactc caaggcttca gccgacttt 180  
 cctgaaaggta aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240  
 gggccctccctt gggactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
 cctccctcaggc cacctccaga tcgacaaggat acccaggatg gaggagaagg aggccctgg 360  
 acccatccag aaggccacgg acagttcca cacagaactc catccccggg tggccttctg 420  
 gatcattaag ctgcccacggc ggaggttcca ccaggatgcc ctggaggccg gccactggct 480  
 cagcgagaag cgacaccggc tgcaggccat ccggatggc ctccgcaagg ggacccacaa 540  
 ggacgtccta gaagagggga ccgagagctc ctcccactcc aggtgttccc cccgaaagac 600  
 ccacttactg tacatcctca ggccctctcg gcagctgtag 640

&lt;210&gt; 14

&lt;211&gt; 211

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 14

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
 50 55 60  
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
 65 70 75 80  
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
 85 90 95  
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Val Pro Arg Met  
 100 105 110  
 Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe  
 115 120 125  
 His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro  
 130 135 140  
 Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser  
 145 150 155 160  
 Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly  
 165 170 175  
 Thr His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser  
 180 185 190  
 Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser  
 195 200 205  
 Arg Gln Leu  
 210

&lt;210&gt; 15

&lt;211&gt; 640

&lt;212&gt; DNA

## SEQLIST.TXT

&lt;213&gt; homo sapien

&lt;400&gt; 15

caccatggga gaaggcctccc cacctgcccc cgcaaggcgg catctgctgg tcctgctgct 60  
 gctcctctct accctggta tccccccgc tgcaagctcct atccatgatg ctgacgccc 120  
 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgacttt 180  
 cctgaaaggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240  
 gggcctccct gggactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300  
 cctctccagc cacctccaga tcgacaaggt acccaggatg gaggagaagg aggccttgg 360  
 acccatccag aaggccacgg acagttcca cacagaactc catccccggg tggcctctg 420  
 gatcatthaag ctgccacggc ggaggtccca ccaggatgcc ctggaggggc gccactggct 480  
 cagcgagaag cgacaccggc tgcaggccat cgggatgga ctccgcaagg ggaccaccaa 540  
 ggacgtccta gaagaggaga ccgagagctc ctcccactcc aggtgtccc cccgaaagac 600  
 ccacttactg tacatcctca ggccctctcg gcagctgttag 640

&lt;210&gt; 16

&lt;211&gt; 211

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 16

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu  
 50 55 60  
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly  
 65 70 75 80  
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly  
 85 90 95  
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Val Pro Arg Met  
 100 105 110  
 Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe  
 115 120 125  
 His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro  
 130 135 140  
 Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser  
 145 150 155 160  
 Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly  
 165 170 175  
 Thr His Lys Asp Val Leu Glu Glu Thr Glu Ser Ser Ser His Ser  
 180 185 190  
 Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser  
 195 200 205  
 Arg Gln Leu  
 210

&lt;210&gt; 17

&lt;211&gt; 499

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 17

caccatggga gaaggcctccc cacctgcccc cgcaaggcgg catctgctgg tcctgctgct 60  
 gctcctctct accctggta tccccccgc tgcaagctcct atccatgatg ctgacgccc 120  
 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgacttt 180  
 cctgaaagta cccaggatgg aggagaagga ggccttggta cccatccaga aggccacgg 240  
 cagcttccac acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggc 300  
 gaggtccccac caggatgccc tggaggggcag ccactggctc agcgagaagc gacaccgc 360

## SEQLIST.TXT

gcaggccatc cgggatggac tccgcaaggg gacccacaag gacgtcctaa aagaggggac 420  
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 gccctctcgg cagctgtag 499

<210> 18  
 <211> 164  
 <212> PRT  
 <213> homo sapien

<400> 18  
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
 1 5 10 15  
 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg  
 50 55 60  
 Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser  
 65 70 75 80  
 Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu  
 85 90 95  
 Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu  
 100 105 110  
 Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys  
 115 120 125  
 Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His  
 130 135 140  
 Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro  
 145 150 155 160  
 Ser Arg Gln Leu

<210> 19  
 <211> 499  
 <212> DNA  
 <213> homo sapien

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 agagagctcc ttgggtctca caggcctcca gaggctactc caaggcttca gccgacttt 180  
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 gcaggccatc cgggatggac tccgcaaggg gacccacaag gacgtcctaa aagaggggac 420  
 cgagagctcc tcccactcca ggctgtcccc ccgaaagacc cacttactgt acatcctcag 480  
 gccctctcgg cagctgtag 499

<210> 20  
 <211> 164  
 <212> PRT  
 <213> homo sapien

<400> 20  
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 1 5 10 15  
 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg

## SEQLIST.TXT

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Phe	His	Thr	Glu	Leu	His	Pro	Arg	Val	Ala	Phe	Trp	Ile	Ile	Lys	Leu
										85					95
Pro	Arg	Arg	Arg	Ser	His	Gln	Asp	Ala	Leu	Glu	Gly	Ser	His	Trp	Leu
										100	105				110
Ser	Glu	Lys	Arg	His	Arg	Leu	Gln	Ala	Ile	Arg	Asp	Gly	Leu	Arg	Lys
										115	120				125
Gly	Thr	His	Lys	Asp	Val	Leu	Lys	Glu	Gly	Thr	Glu	Ser	Ser	Ser	His
										130	135				140
Ser	Arg	Leu	Ser	Pro	Arg	Lys	Thr	His	Leu	Leu	Tyr	Ile	Leu	Arg	Pro
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Ser	Arg	Gln	Leu												160

&lt;210&gt; 21

&lt;211&gt; 499

&lt;212&gt; DNA

&lt;213&gt; homo sapien

&lt;400&gt; 21

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agagagctcc	ttgggtctca	caggcctcca	gagcctactc	caaggcttca	gccgactttt	180
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gaggtcccac	caggatgccc	tggagggcag	ccactggctc	agcgagaagc	gacaccgcct	360
gcagggccatc	cgggatggac	tccgcaaggg	gaccacaaag	gacgtcttag	aagaggggac	420
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&lt;210&gt; 22

&lt;211&gt; 164

&lt;212&gt; PRT

&lt;213&gt; homo sapien

&lt;400&gt; 22

Met	Gly	Glu	Ala	Ser	Pro	Pro	Ala	Pro	Ala	Arg	Arg	His	Leu	Leu	Val
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Leu	Leu	Leu	Leu	Leu	Ser	Thr	Leu	Val	Ile	Pro	Ser	Ala	Ala	Ala	Pro
										20			25		30
Ile	His	Asp	Ala	Asp	Ala	Gln	Glu	Ser	Ser	Leu	Gly	Leu	Thr	Gly	Leu
										35			40		45
Gln	Ser	Leu	Leu	Gln	Gly	Phe	Ser	Arg	Leu	Phe	Leu	Lys	Val	Pro	Arg
										50			55		60
Met	Glu	Glu	Glu	Ala	Leu	Val	Pro	Ile	Gln	Lys	Ala	Thr	Asp	Ser	
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Phe	His	Thr	Glu	Leu	His	Pro	Arg	Val	Ala	Phe	Trp	Ile	Ile	Lys	Leu
										85			90		95
Pro	Arg	Arg	Arg	Ser	His	Gln	Asp	Ala	Leu	Glu	Gly	Ser	His	Trp	Leu
										100			105		110
Ser	Glu	Lys	Arg	His	Arg	Leu	Gln	Ala	Ile	Arg	Asp	Gly	Leu	Arg	Lys
										115			120		125
Gly	Thr	His	Lys	Asp	Val	Leu	Glu	Glu	Gly	Thr	Glu	Ser	Ser	Ser	His
										130			135		140
Ser	Arg	Leu	Ser	Pro	Arg	Lys	Thr	His	Leu	Leu	Tyr	Ile	Leu	Arg	Pro
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Ser	Arg	Gln	Leu												

## SEQLIST.TXT

<210> 23  
 <211> 499  
 <212> DNA  
 <213> homo sapien

<400> 23  
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 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgacttt 180  
 cctgaaagta cccaggatgg aggagaagga ggccctggta cccatccaga aggcccacgga 240  
 cagcttccac acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggcg 300  
 gaggtcccac caggatgccc tggagggcag ccactggctc aegcagaagc gacaccgcct 360  
 gcaggccatc cgggatggac tccgcaaggg gaccacaaag gacgtcctaa aagaggggac 420  
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 gcccctctcg cagctgttag 499

<210> 24  
 <211> 164  
 <212> PRT  
 <213> homo sapien

<400> 24  
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 Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro  
 20 25 30  
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
 35 40 45  
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg  
 50 55 60  
 Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser  
 65 70 75 80  
 Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu  
 85 90 95  
 Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu  
 100 105 110  
 Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys  
 115 120 125  
 Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His  
 130 135 140  
 Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro  
 145 150 155 160  
 Ser Arg Gln Leu

<210> 25  
 <211> 499  
 <212> DNA  
 <213> homo sapien

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 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgacttt 180  
 cctgaaagta cccaggatgg aggagaagga ggccctggta cccatccaga aggcccacgga 240  
 cagcttccac acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggcg 300  
 gaggtcccac caggatgccc tggagggcag ccactggctc aegcagaagc gacaccgcct 360  
 gcaggccatc cgggatggac tccgcaaggg gaccacaaag gacgtcctaa aagaggggac 420  
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<210> 26

SEQLIST.TXT

<211> 164  
<212> PRT  
<213> homo sapien

<400> 26  
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val  
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Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro  
20 25 30  
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu  
35 40 45  
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg  
50 55 60  
Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser  
65 70 75 80  
Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu  
85 90 95  
Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu  
100 105 110  
Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys  
115 120 125  
Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His  
130 135 140  
Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro  
145 150 155 160  
Ser Arg Gln Leu